## ES prok arch Differential expression HvN

All ✓ records per page	Search all columns:			
		From to	From	From
ID	lman.	le=EC	p-	Adjust
k141_936623_8_Belongs_to_the_sodium_solute_symport	Image	-0.791	1.10e-04	1.69e-0
K141_000020_0_bclosligo_lo_tile_bouldin_boulde_symport	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_936623_8_Belongs_to	-0.701	1.100-04	1.000-0
k141_3636_26_TPR_repeat	6	0.929	7.09e-04	5.75e-C
k141_3511_15		1.460	5.05e-04	4.58e-C
k141_787459_1_Stabilizes_TBP_binding_to_an_archaea	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_787459_1_Stabilizes  o	1.840	3.82e-04	3.69e-C
k141_229285_13_copper_ion_binding	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_229285_13_copper_io	1.850	1.88e-05	7.25e-C
k141_229285_9_DnaJ_molecular_chaperone_homology_do	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_229285_9_DnaJ_mole	1.970	2.38e-04	2.83e-C
k141_3511_26	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_3511_26pdf)	2.090	2.19e-04	2.81e-C
k141_3636_2	o (figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_3636_2pdf)	2.260	2.90e-04	2.99e-C
k141_812962_2_COG0589_Universal_stress_protein_Usp	6	2.340	5.34e-05	1.37e-C
k141_3511_10_amino_acid	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_3511_10_amino_acid	2.560	6.35e-05	1.40e-C
k141_229285_6_Belongs_to_the_LDH_MDH_superfamily	o 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	2.590	6.99e-04	5.75e-C
k141_229285_12_Belongs_to_the_P_IIprotein_family	♦ ♦ ⑤ ○ ○ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦ ♦	2.750	7.90e-04	6.09e-C
k141_3511_11	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_3511_11pdf)	2.770	1.29e-04	1.80e-C
k141_3511_14_Belongs_to_the_small_heat_shock_prote	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_3511_14_Belongs_to	2.820	7.30e-05	1.41e-C
k141_229285_5_CBS_domain	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_229285_5_CBS_dom	2.970	9.19e-05	1.58e-C
k141_24439_4_CBS_domain	6	3.150	2.66e-04	2.93e-C
k141_3511_13_Cell_division_protein_48CDC48dom	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_3511_13_Cell_division	3.150	1.14e-05	5.85e-C
k141_229285_8	o old on the control of the control	3.180	7.06e-09	1.09e-C
k141_696673_1_Belongs_to_the_P_IIprotein_family	(figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_696673_1_Belongs_to	3.490	4.06e-05	1.25e-C
k141_24439_3_transcriptional_regulators	6-023 9-6 (figuresES_prok_arch_RNAseq_analysis_with_DESeq2_long_names_HvN/boxplot.k141_24439_3_transcription 0 60 89-0	4.290	2.45e-06	1.89e-C
Showing 1 to 20 of 20 entries	$\leftarrow$ Previous Next $\rightarrow$			